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TITLE OF INVENTION: Medical Uses of These Substances
FILE REFERENCE: 010830-117
CURRENT APPLICATION NUMBER: US/09/840,243B
CURRENT FILING DATE: 2001-04-24
PRIOR APPLICATION NUMBER: EP 98120085.0
PRIOR FILING DATE: 1998-10-24
NUMBER OF SEQ ID NOS: 22
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 12
LENGTH: 260
TYPE: PRT
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   APPLICANT: MASTERNAK, Krzysztof

APPLICANT: REITH, Walter

APPLICANT: MACH, Bernard

TITLE OF INVENTION: New Transcription Factor of MHC Clas

TITLE OF INVENTION: Capable of Inhibiting This New Tran

TITLE OF INVENTION: Medical Uses of These Substances

FILE REFERENCE: 010830-117

CURRENT APPLICATION NUMBER: US/09/840,243B

CURRENT FILING DATE: 2001-04-24

PRIOR APPLICATION NUMBER: EP 98120085.0

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PRIOR FILING DATE: 1998-10-24
NUMBER OF SEQ ID NOS: 22
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 18
LENGTH: 220
TYPE: PRT
ORGANISM: Homo sapiens
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APPLICANT: REITH, Walter
APPLICANT: MACH, Bernard
TITLE OF INVENTION: New Transcription Factor of MHC Class II Genes, Substitute Of Invention: Capable of Inhibiting This New Transcription Factor TITLE OF INVENTION: Medical Uses of These Substances
FILE REFERENCE: 010830-117
                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: CURRENT FILING DATE: 2001-
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Best Local S
Matches 112
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Patent No.
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APPLICANT: MASTERNAK,
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APPLICANT:
TITLE OF IN
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CURRENT APPLICATION NUMBER: US/09/840,243B
CURRENT FILING DATE: 2001-04-24
PRIOR APPLICATION NUMBER: EP 98120085.0
PRIOR FILING DATE: 1998-10-24
NUMBER OF SEQ ID NOS: 22
PRIOR PRIOR PRIOR PRIOR
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                                              APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED ITTLE OF INVENTION: GENE EXPRESSION ANALYTICE REFERENCE: Aeomica-X-1
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
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TYPE: PRT
ORGANISM: Murinae
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FILING DATE: 2000-02-04
APPLICATION NUMBER: US 60/207,
FILING DATE: 2000-05-26
APPLICATION NUMBER: US 09/632,
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REITH, war-
MACH, Bernard
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larity 58.9%;
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Transcription
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OR APPLICATION NUMBER: PCT/US01/00666

OR APPLICATION NUMBER: PCT/US01/00666

OR APPLICATION NUMBER: PCT/US01/00667

OR APPLICATION NUMBER: PCT/US01/00667

OR APPLICATION NUMBER: PCT/US01/00664

OR APPLICATION NUMBER: PCT/US01/00664

OR APPLICATION NUMBER: PCT/US01/00669

OR APPLICATION NUMBER: PCT/US01/00669

OR APPLICATION NUMBER: PCT/US01/00665

OR APPLICATION NUMBER: PCT/US01/00668

OR FILING DATE: 2001-01-30

OR APPLICATION NUMBER: PCT/US01/00663

OR APPLICATION NUMBER: PCT/US01/00663

OR APPLICATION NUMBER: PCT/US01/00661

OR APPLICATION NUMBER: PCT/US01/00661

OR APPLICATION NUMBER: US 60/234,687

OR APPLICATION NUMBER: US 60/234,687

OR APPLICATION NUMBER: US 09/608,408

OR APPLICATION NUMBER: US 09/608,408

OR APPLICATION NUMBER: US 09/774,203

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INFORMATION:
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EXPRESSED IN HELA, SIGNAL = 1.9

EXPRESSED IN LUNG, SIGNAL = 2.5

EXPRESSED IN FETAL LIVER, SIGNAL = 1.7

EST_HUMAN HIT: BE905633.1, EVALUE 3.00e-22

SWISSPROT HIT: O14593, EVALUE 2.00e-23
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Pred. No. 1e-
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US-09-835-788A-17

y Sequence 17, Application US/09835788A

patent No. US20020077458A1

patent No. US20020077458A1

GENERAL INFORMATION:

APPLICANT: Ni et al.

TITLE OF INVENTION: Death Domain-Containing Receptor Polynucleotides,

TITLE OF INVENTION: Antibodies

FILE REFERENCE: PT018P1

CURRENT APPLICATION NUMBER: US/09/835,788A

CURRENT FILING DATE: 2001-04-17

PRIOR FILING DATE: 2000-10-17

PRIOR APPLICATION NUMBER: PCT/US00/28666

PRIOR APPLICATION NUMBER: 60/159,585

PRIOR APPLICATION NUMBER: 60/159,585

PRIOR FILING DATE: 1999-10-18

PRIOR FILING DATE: 1999-11-24

NUMBER OF SEQ ID NOS: 24
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Polypeptides,

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SEQ ID NO 12
LENGTH: 740
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Best Local S
Matches 58
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CURRENT FILING DATE: 2001-04-17
PRIOR APPLICATION NUMBER: PCT/US00/28666
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/159,585
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PRIOR APPLICATION NUMBER: 6
PRIOR FILING DATE: 1999-11-
NUMBER OF SEQ ID NOS: 24
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TYPE: PRT
ORGANISM: Homo
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ORGANISM: Homo
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TITLE OF INVENTION
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                                     LLLERDVD---INIYDWNGGTPLLYAVRGNHVKCVEALLARGADLTTEADS--GYTPMDL
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No. 9.3e-10;
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Patent No

INFORM

ATION:

No.

US 20

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APPLICANT:

APPLICANT:

de Lang Smith,

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METHODS

TITLE OF INVENTION: A PROTEIN THAT BINDS TITLE OF INVENTION: OF USE THEREOF NUMBER OF SEQUENCES: 12

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RESULT 10
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FILING DATE: 2000-01-31
APPLICATION NUMBER: 60/180,628
FILING DATE: 2000-02-04
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Application US/09841835

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RESULT 11
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CURRENT APPLICATION
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                                                                                                                  GGTPLLYAVRGNHVKCVEALLARGADLTTEADSGYTPMDLAVALGYRKVQQVIENH
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CHARACTERISTICS:
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: IBM PC compatible
G SYSTEM: PC-DOS/MS-DOS
: PatentIn Release #1.0,
PLICATION DATA:
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larity 24.3%;
Conservative 3
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Pred. No. 9e-09;
36; Mismatches 1
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US-09-972-115A-8
; Sequence 8, Application
; Publication No. US200300; GENERAL INFORMATION:
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Best Local S
Matches 72
                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: proto
3-09-841-835-10
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APPLICATION NUMBER: US/no/
FILING DATE:
CLASSIFTCT
APPLICANT:
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APPLICANT:
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INFORMATION FOR SEQ
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ZIP: 07601
COMPUTER READABLE I
MEDIUM TYPE: F1c
COMPUTER: IBM PC
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TITLE OF 1
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NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600-1-230
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800
TELEFAX: 201-343-1684
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CORRESPONDENCE ADDRESS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
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LENGTH: 949 amino acids
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Pred. No. 1.5e-08;
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Patent No
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ORGANISM: Homo
-09-972-115A-8
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CURRENT APPLICATION NUMBER: US/09/972,1:
CURRENT FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 60/128,577
PRIOR FILING DATE: 2000-04-10
PRIOR APPLICATION NUMBER: US 60/129,123
PRIOR FILING DATE: 1999-04-13
CURRENT A...

CURRENT A...

APPLICATION NUMBER:

CLASSIFICATION:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/196,

FILING DATE:

ATTORNEY/AGENT INFORMATION:

NAME: Jackson Esq., David

NAME: Jackson NUMBER: 26,7/
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COMPUTER READAB
MEDIUM TYPE:
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TITLE OF IN
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CORRESPONDEN
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OPERATING SYSTEM: PC-D
SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
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CITY: Ha
STATE: N
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). US20020076795A1
INFORMATION:
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                                                                                                                                                                    PE: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0,
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Smith, Susan
ZENTION: A PROTEIN THAT
ZENTION: OF USE THEREOF
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ATION NUMBER: US/09/972,115A
DATE: 2001-10-05
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Pred. No. 2.3e-08;
36; Mismatches 112;
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                                                                                                                                                                                                        TITLE OF INVENTION: Death Domain-Containing Re
TITLE OF INVENTION: Antibodies
FILE REFERENCE: PT018P1
CURRENT APPLICATION NUMBER: US/09/835,788A
CURRENT FILING DATE: 2001-04-17
PRIOR APPLICATION NUMBER: PCT/US00/28666
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/159,585
PRIOR APPLICATION NUMBER: 60/159,585
PRIOR APPLICATION NUMBER: 60/167,246
PRIOR FILING DATE: 1999-10-18
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 24
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                                                                                  Best
                                                                 Matches
                                                                                                                                                                  SEQ ID NO 18
LENGTH: 285
                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18, Application US/09835788A Patent No. US20020077458A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Ni et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
HYPOTHETICAL:
09-841-835-2
                                                                                                                                   TYPE: PRT
ORGANISM: Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 133521
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 1327 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 60
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800
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TOPOLOGY: 111
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KGHLDMVRFLLEAGADQEHKTDEM-HTALMEACMDGHVEVARLLLDSGAQVNMPADSFES
                            QGELDQLKEHLRKGDNLVNKPDERGFTPLIWASAFGEIETVRFLLEWGADPHILAKERES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HILAKERESALSLAS----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KSTPLHLAAGYNRVRIVQLLLQHGADVHAKDKGGLVPLHNACSYGHYEVTELLLKH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGTPLLYAVRGNHVKCVEALLARGADLTTEADSGYTPMDLAVALGYRKVQQVIENH
                                                                                                                            788A-18
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                                                                 48;
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L Similarity 24.3%;
72; Conservative
                                                                            Similarity
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                                                          Score 185; DB 1
Pred. No. 4e-09;
0; Mismatches
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Pred. No. 2.3e-08;
5; Mismatches 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 --EW-
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                                                                                      DB 10;
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                                                                                       Length
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US-09-964-899-43
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                                                                                                                                                                                                                                                                                                                    SOFTWARE: Fa
SEQ ID NO 43
LENGTH: 172
                                                                                                                                                                                                      Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent
                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo
09-964-899-43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Cohen, Dalia et al.
TITLE OF INVENTION: Identification of Genes Involved in
TITLE OF INVENTION: Alzheimer's Disease Using Drosophila Melanogaster
FILE REFERENCE: 4-31612 A
  130
                                                                                                      158
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LATEDGFTPLAVALQQGHDQVVSLLLEN 157
                            TEADSGYTPMDLAVALGY-RKVQQVIEN 244
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                                                                                                                                                             AAQGELDQLKEHLRKGDNLVNKPDERGFTPLIWASAFGEIETVRFLLEWGADPHILAKER 157
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3. US20020174446A1
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                                                                                                                                                                                             13.8%; Score 185; DB 9; Length 1:
larity 30.4%; Pred. No. 5.2e-08;
Conservative 36; Mismatches 65; Indels
                                                             LAGQAEVVKVLVTNGANVNAQSQNGFTPLYMAAQENHLEVVKFLLDNGASQS
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